

ESA–MOST Dragon 4 Cooperation

ADVANCED LAND REMOTE SENSING INTERNATIONAL TRAINING COURSE

“龙计划4”高级陆地遥感国际培训班

Rice mapping with polarimetric SAR Data using Sentinel-1 Toolbox

Objectives:

How to process RADARSAT-2 polarimetric products;

Derive Polarimetric decompositions for rice identification;

Rice mapping using unsupervised polarimetric classification and supervised wishart classification.

Data:

The sample data used in this exercise was a subset from one RADARSAT-2 Fine quad-pol image acquired on June 27, 2012.


Pass	Ascending
Product	Single Look Complex (SLC)
Incidence angle mid swath	38.89
Resolution	8.1*7.6 (R*A)
Location	Jiangsu, China

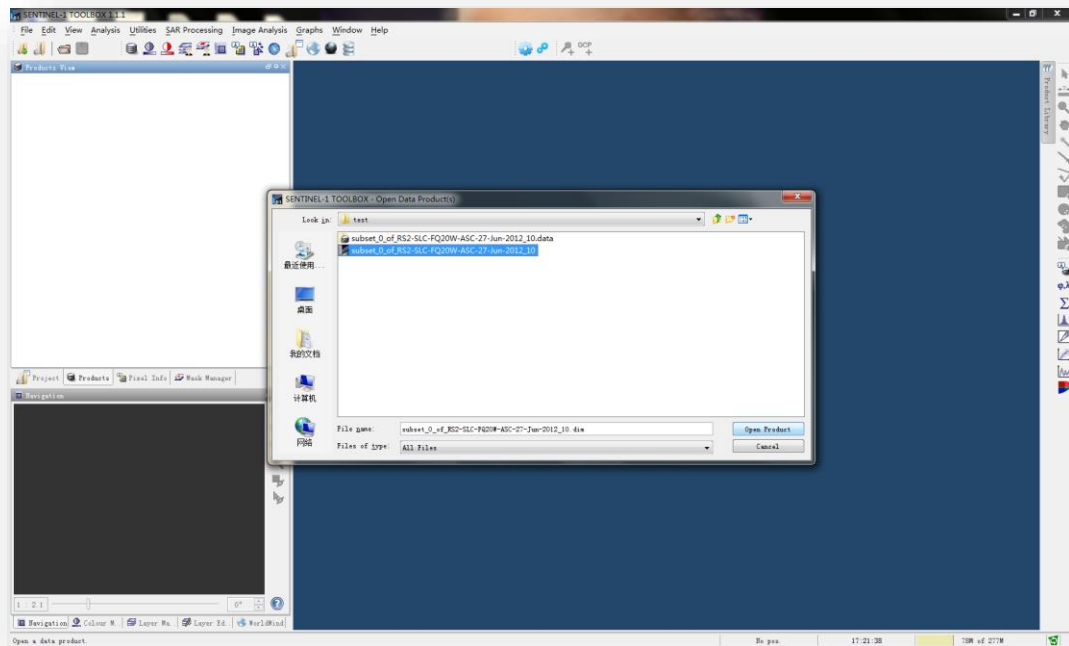
Software:

The Sentinel-1 Toolbox and SNAP, open source software for scientific learning, research and exploitation of the large archives of ESA Sentinel and heritage missions.

Tutorial:

Open the Data:

Use the  **Open Product button** in the top toolbar and browse for the location of the sample data.

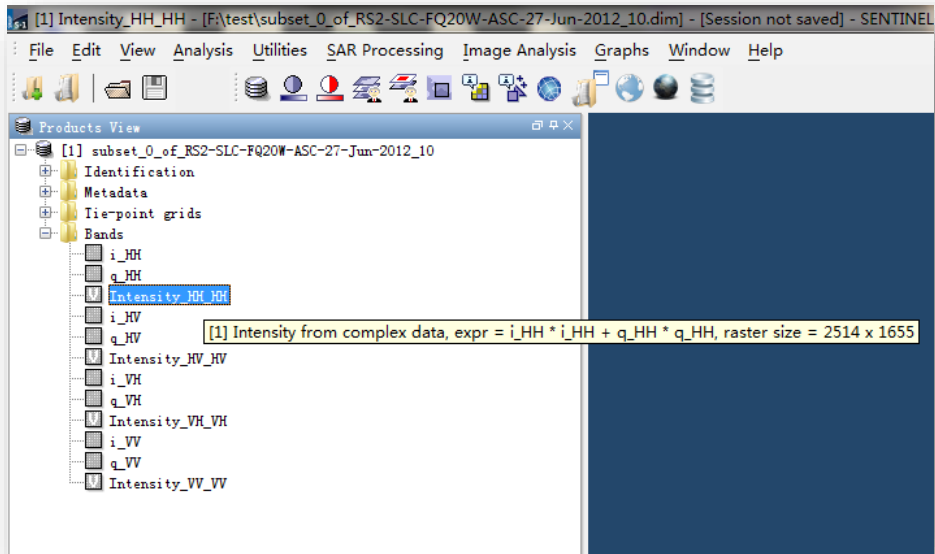


View the product:

In the Products View you will see the opened product. Within the product bands you will see four polarizations:

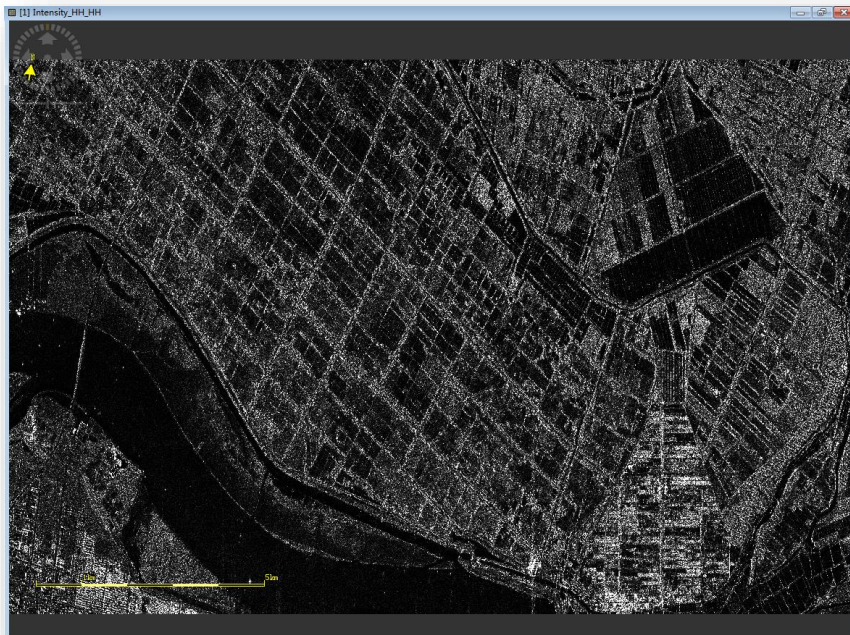
- HH
- HV
- VH
- VV

For each polarization, there will be the complex data **i and q bands** and virtual bands for **intensity**.



View a band:

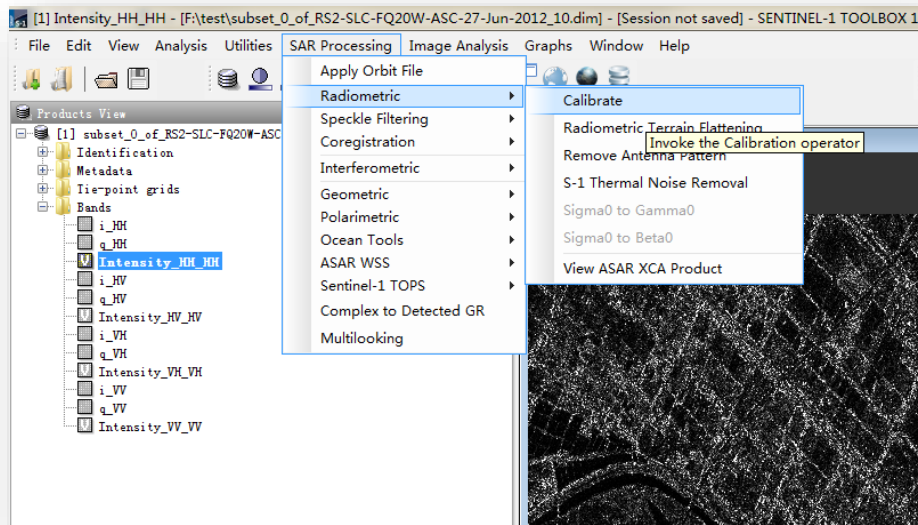
To view the HH band, double-click on the Intensity_HH band. Zoom in using the mouse wheel and pan by clicking and dragging the left mouse button.



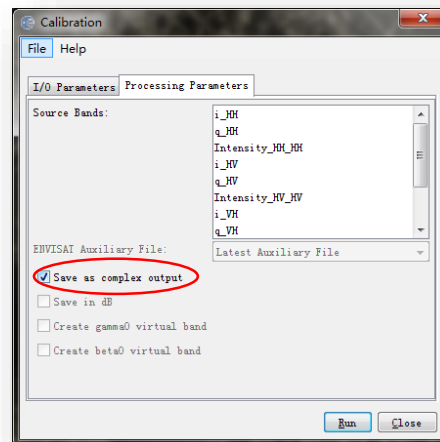
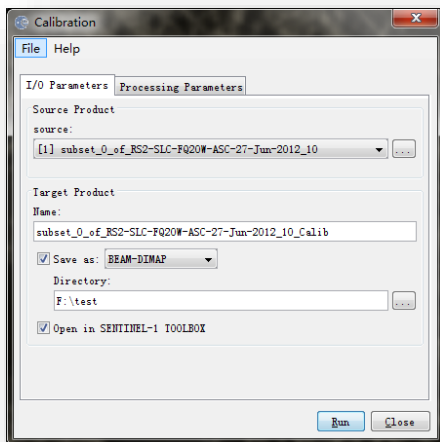
Calibrating the Data

To properly work with the SAR data, the data should first be calibrated. Calibration radiometrically corrects a SAR image so that the pixel values truly represent the radar backscatter of the reflecting surface. Calibration is essential for quantitative use of SAR data.

From the **SAR Processing** menu, go to the **Radiometric** menu and select **Calibrate**.



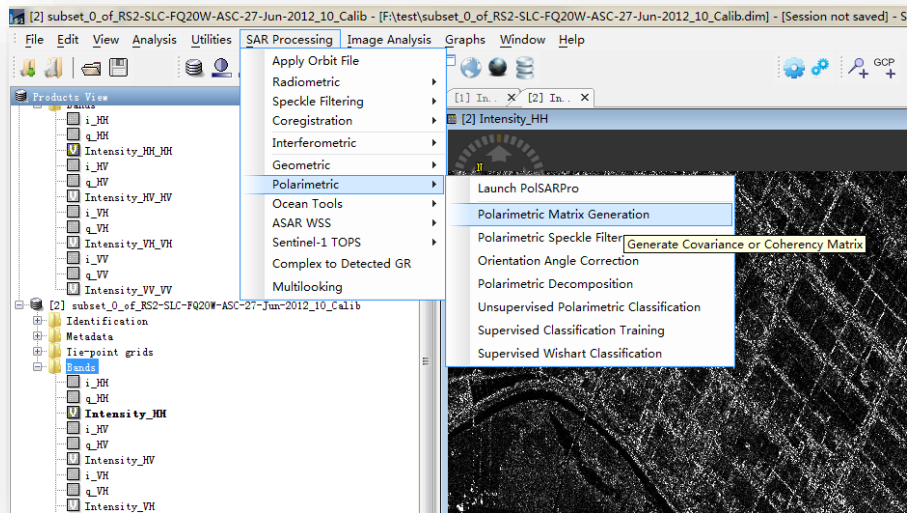
For polarimetric processing the data must be **complex**. In the Processing Parameters tab, check mark the **Save as a complex output**.



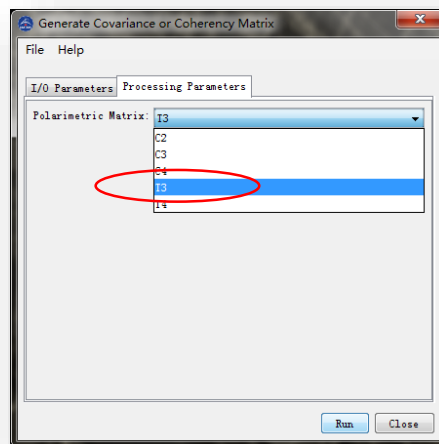
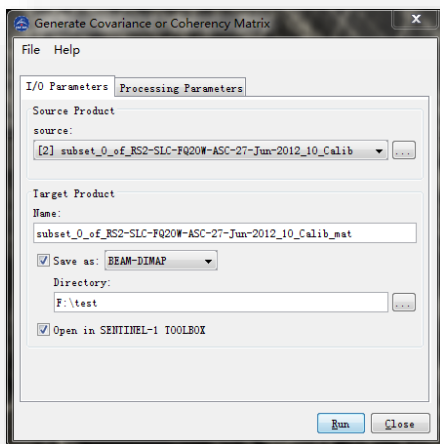
Polarimetric Matrix Generation

All the polarimetric tools work with either Coherency or Covariance matrices as input.

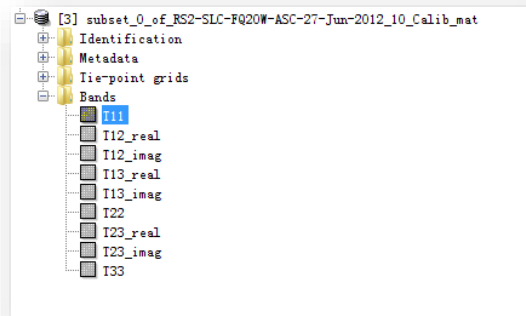
From the **SAR Processing** menu, go to the **Polarimetric** menu and select **Polarimetric Matrix Generation**.



In the **Processing Parameters** tab, select a **T3** matrix to convert the Quad Pol product into a Coherency matrix T3. Press **Run** to begin processing.



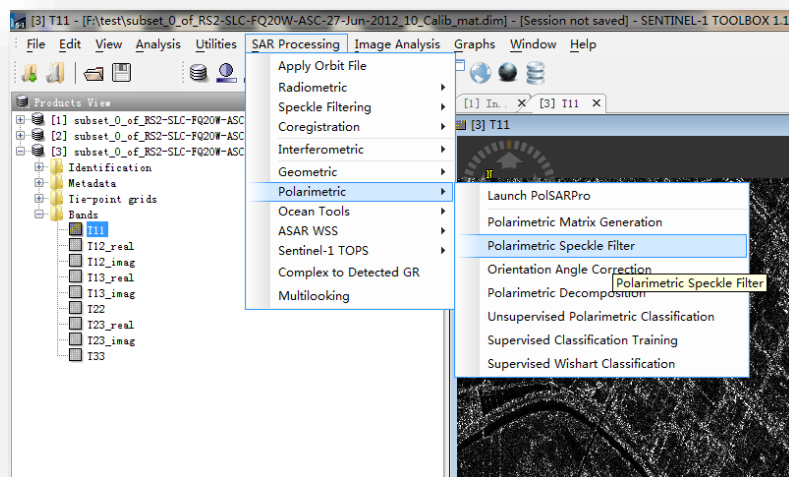
When the processing completes, a new product will be added to the Products View. You will notice the new bands produced correspond to the elements of the T3 matrix.



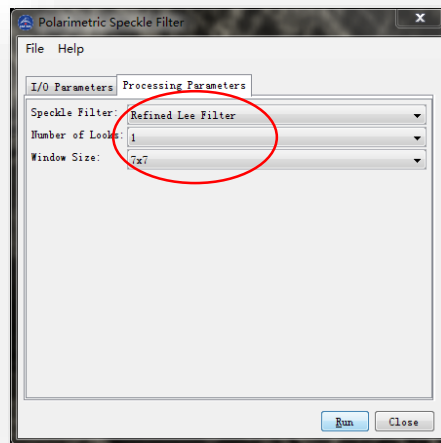
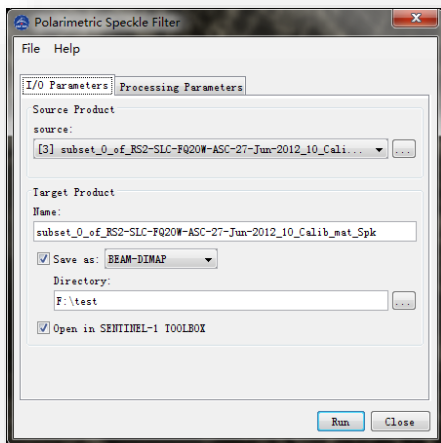
Polarimetric Speckle Filtering

To clean up some of the speckle inherent in SAR images, you can apply a speckle filter. For full polarimetric data, speckle filters must take advantage of all bands and preserve the complex information.

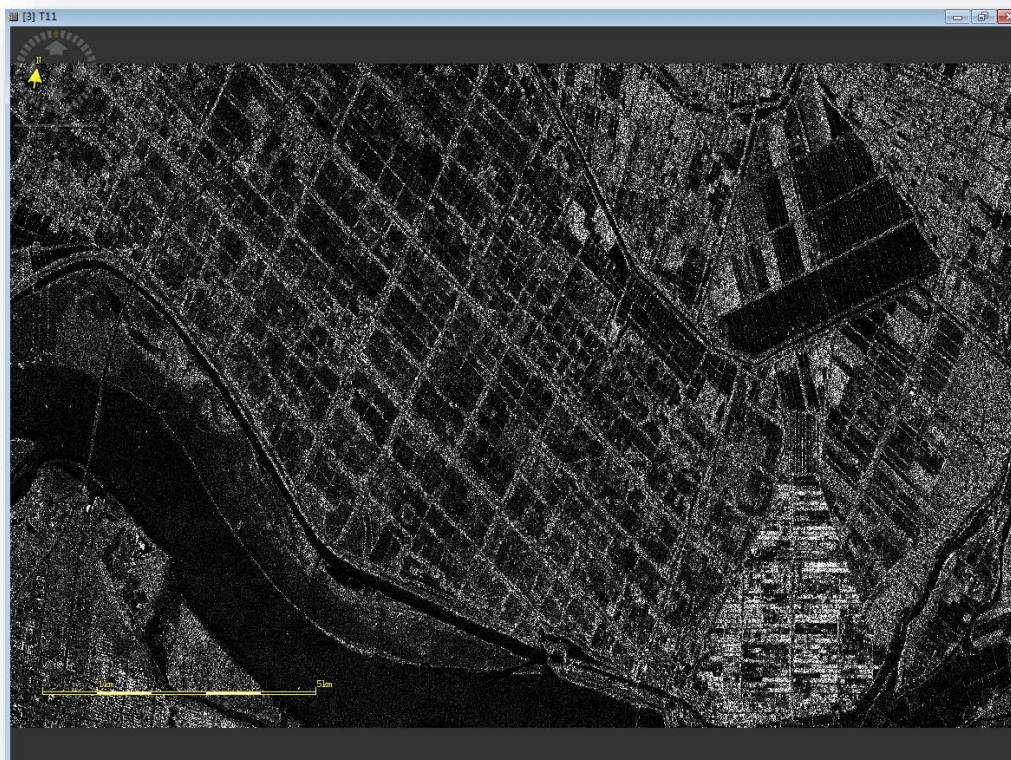
Select **Polarimetric Speckle Filter** from the **Polarimetric** menu.



In the **Processing Parameters** tab, select the **Refined Lee** speckle filter, window size **7x7**. Press **Run** to begin processing.



Open the T11 band in both the T3 product and in new speckle filtered T3 product to compare before and after images. The resulting image will have less speckle but also appear more blurred.

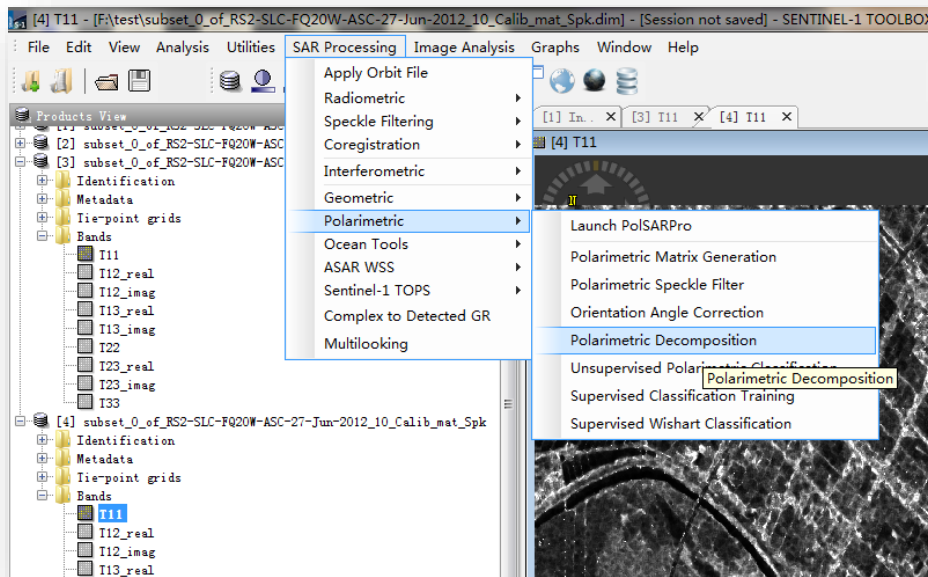




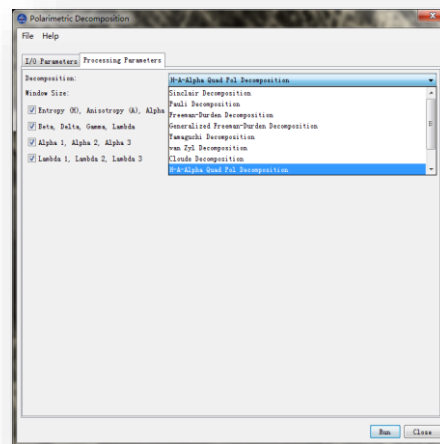
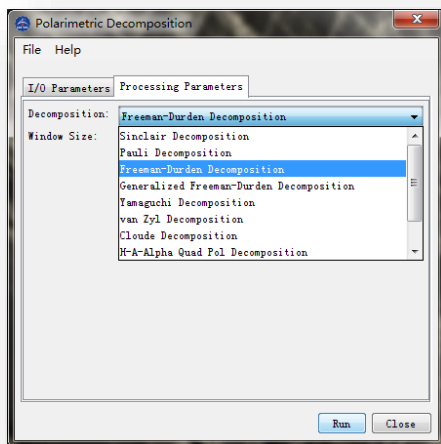
Polarimetric Decompositions

Polarimetric decompositions allow the separation of different scattering contributions and can be used to extract information about the scattering process.

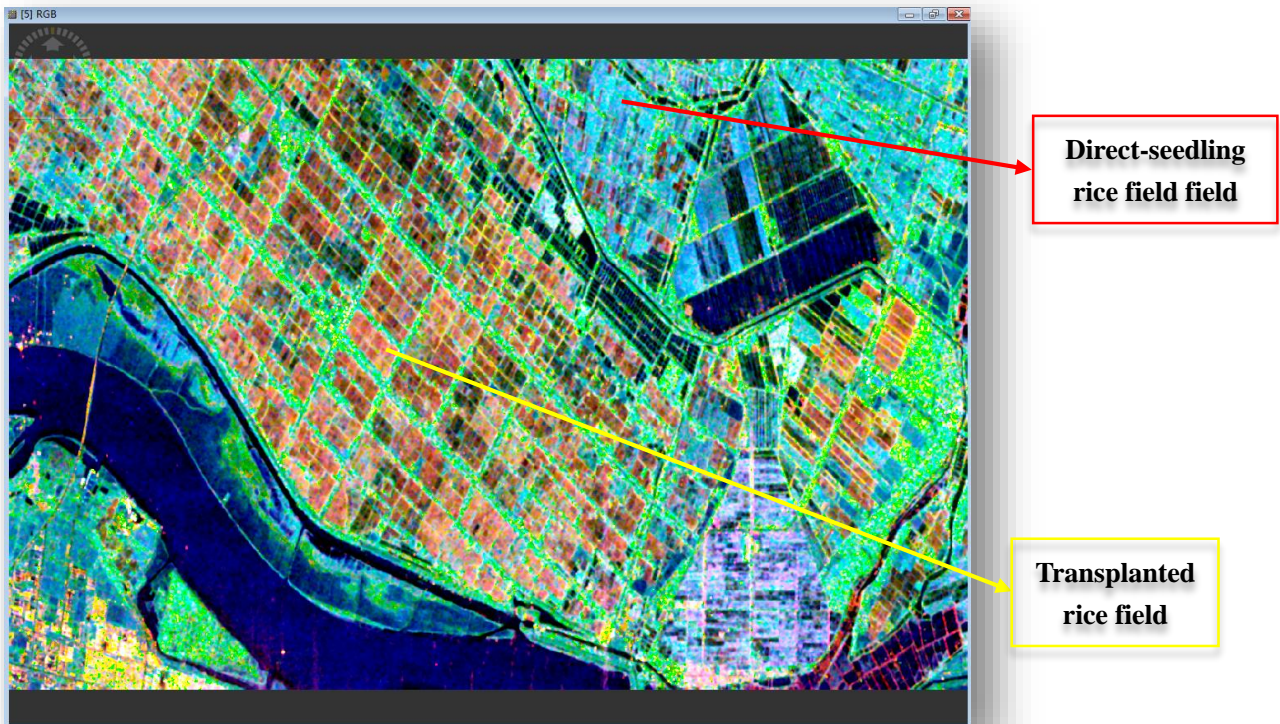
From the **SAR Processing** menu, go to the **Polarimetric** menu and select **Polarimetric Decomposition**.



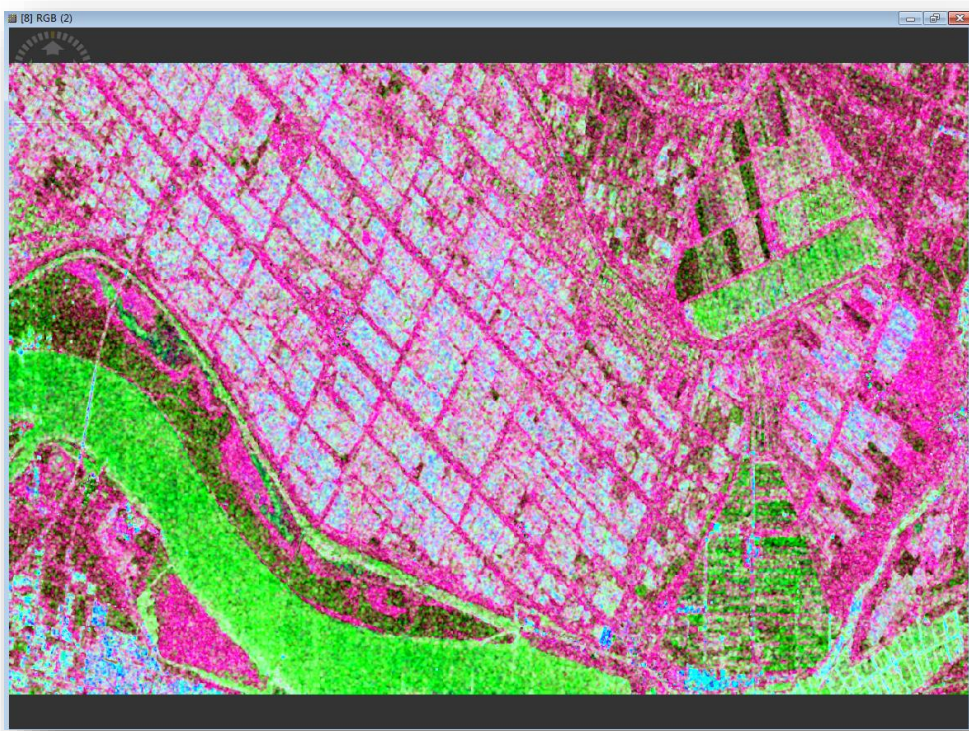
In the **Processing Parameters** tab, select the **Freeman-Durden Decomposition** and **H-A-Alpha** decomposition, window size are both **5x5**. Press **Run** to begin processing.



When the processing completes, you can view all three bands in an RGB colour view by right-clicking on the product name and selecting **Open RGB Image View** from the popup menu.



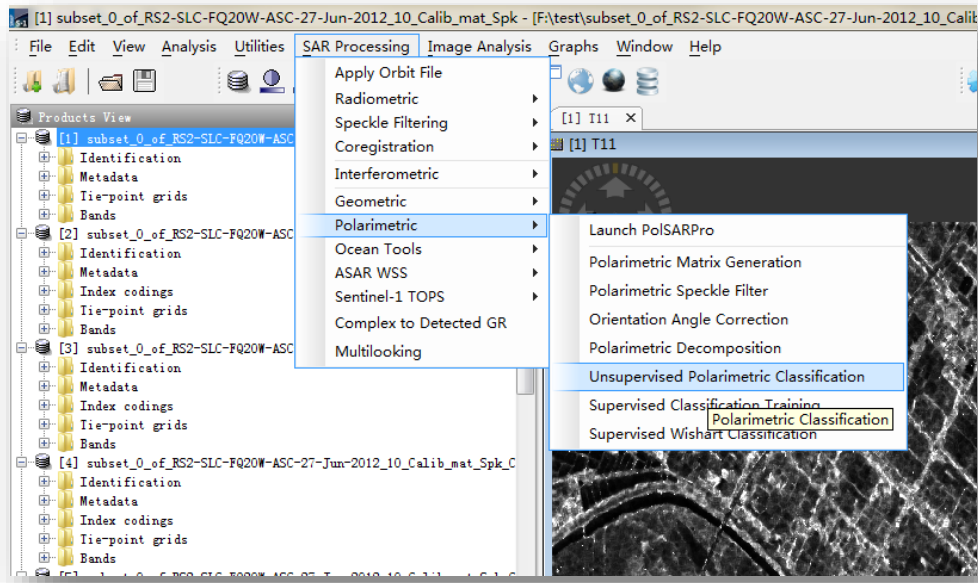
Freeman RGB (R=double bounce G=volume B=surface)



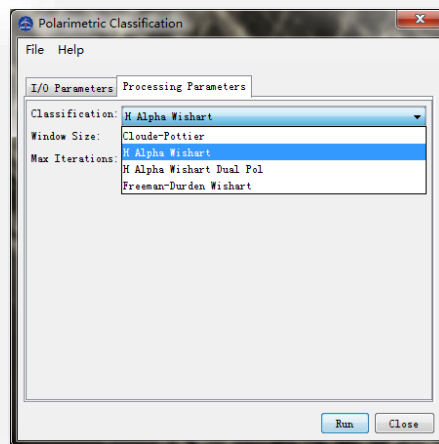
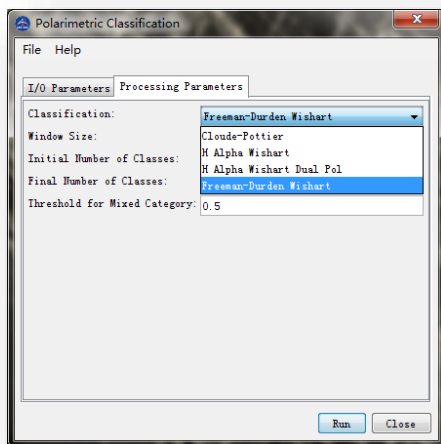
Cloude RGB (R=entropy G=Anisotropy B=alpha)

Unsupervised polarimetric classification

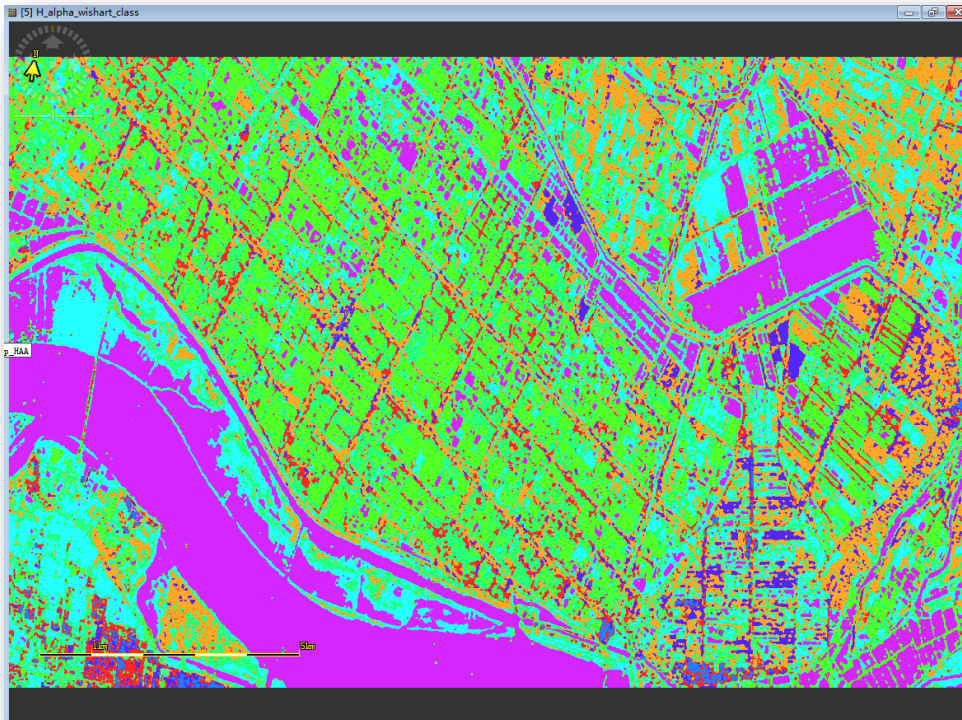
From the **SAR Processing** menu, go to the **Polarimetric** menu and select **Unsupervised polarimetric classification**.



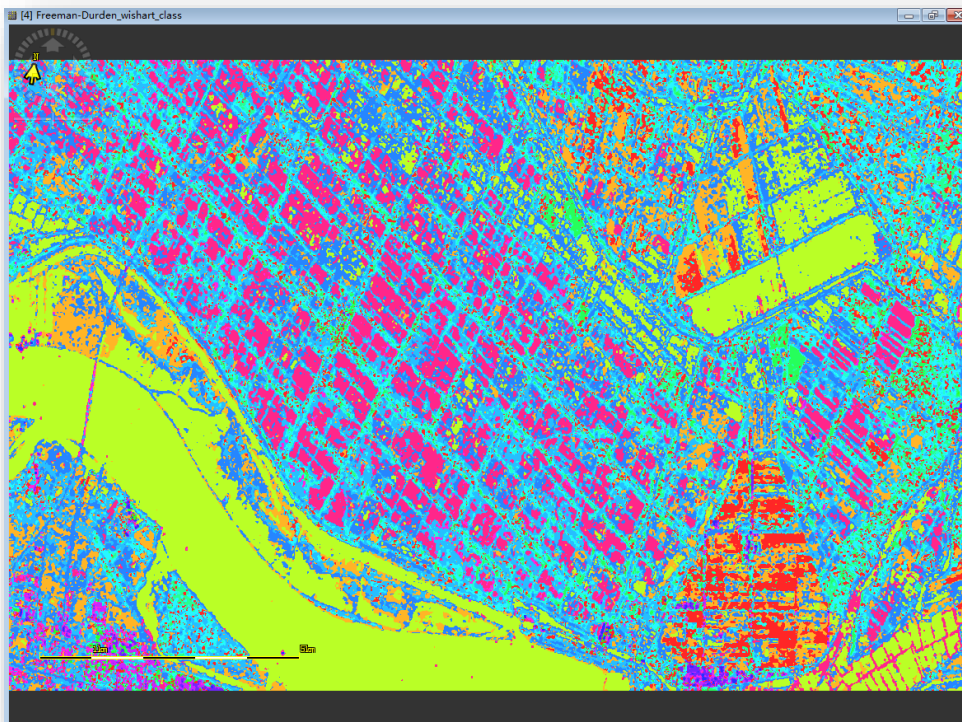
In the **Processing Parameters** tab, select the **H-Alpha Wishart** and **Freeman-Durden Wishart**. Press **Run** to begin processing.



Double-click on the class results to open it.



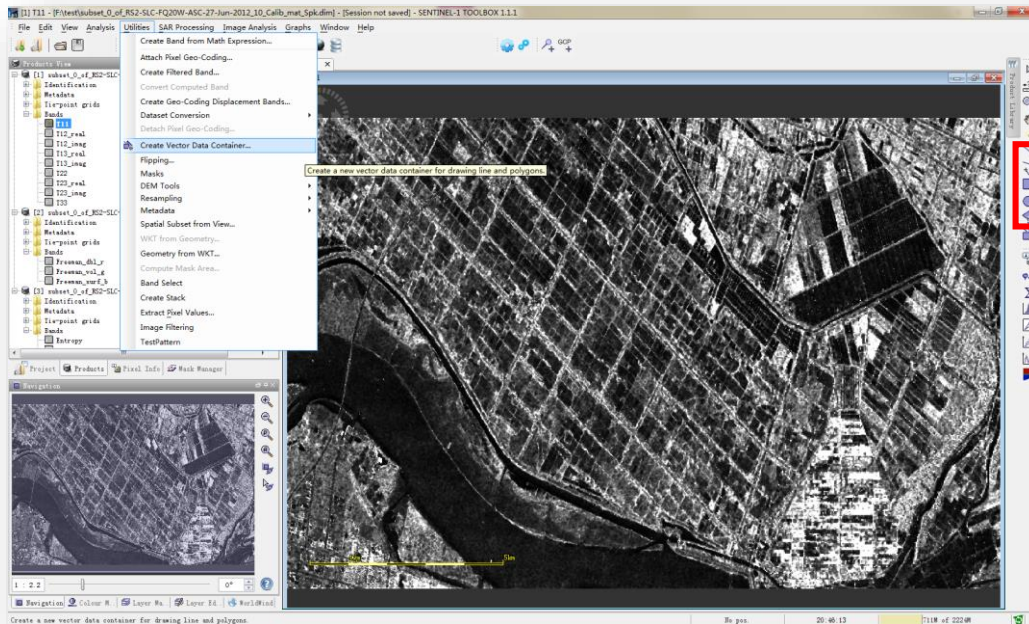
H-Alpha Wishart



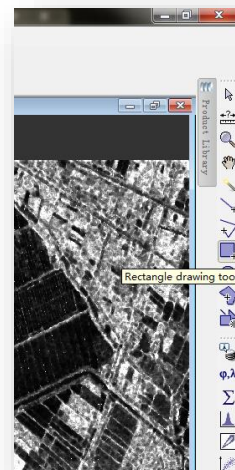
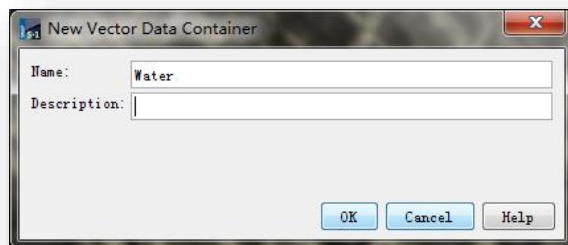
Freeman Wishart

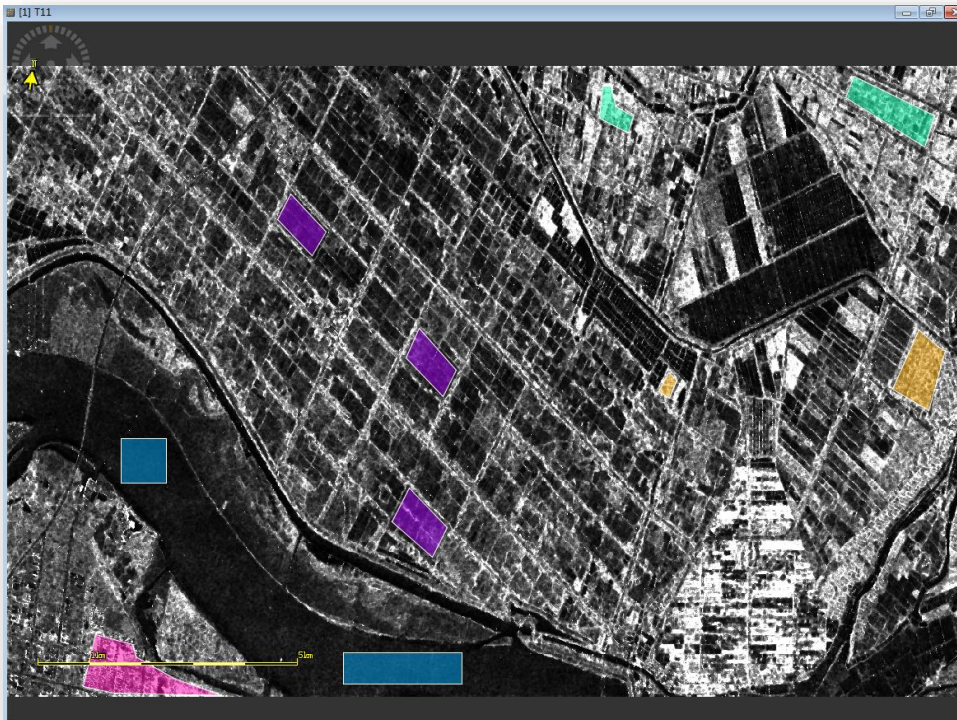
Supervised polarimetric classification

Select areas as training data sets using the **Create a new geometry container** and other drawing tools on the right hand side of the tool box.

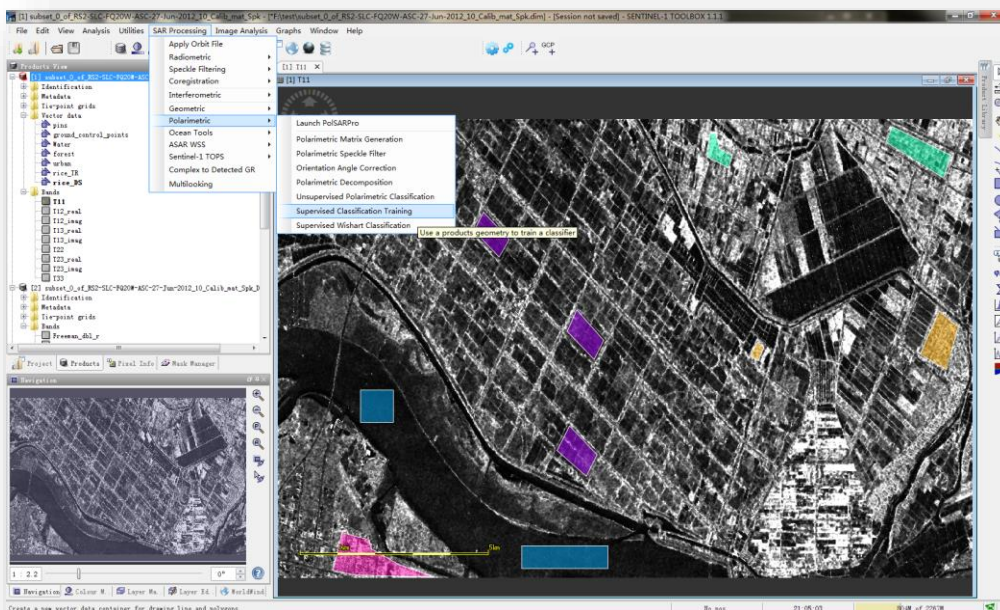


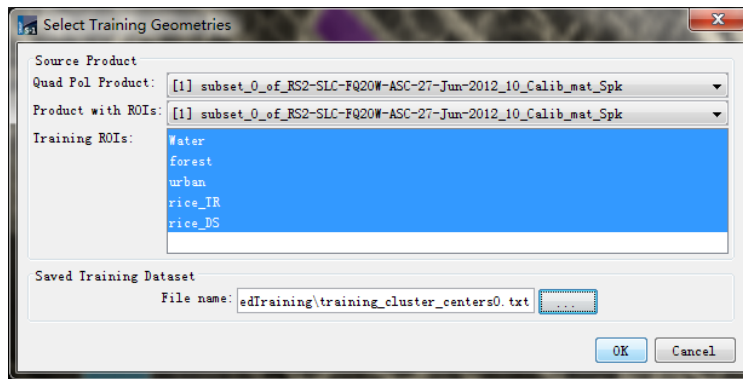
Create a new geometry container, name it and draw it in the image using the drawing tools.



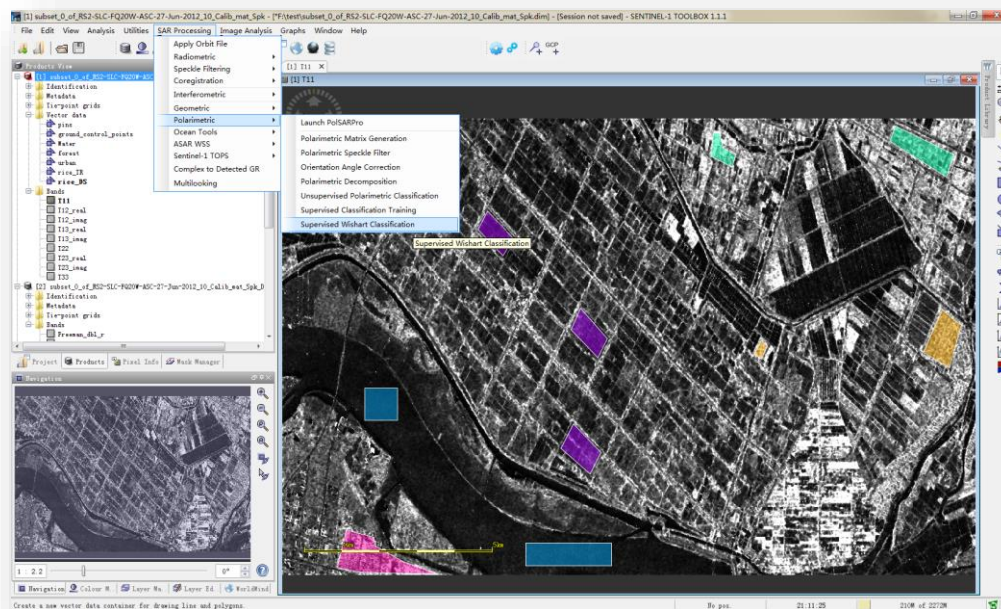


Select **Supervised Classification Training** from the **Polarimetric** menu, then highlight the training geometries and click on "OK" to start the training. The center for the coherency matrices of the pixels in each user identified class is computed and save in a text file in user specified directory.

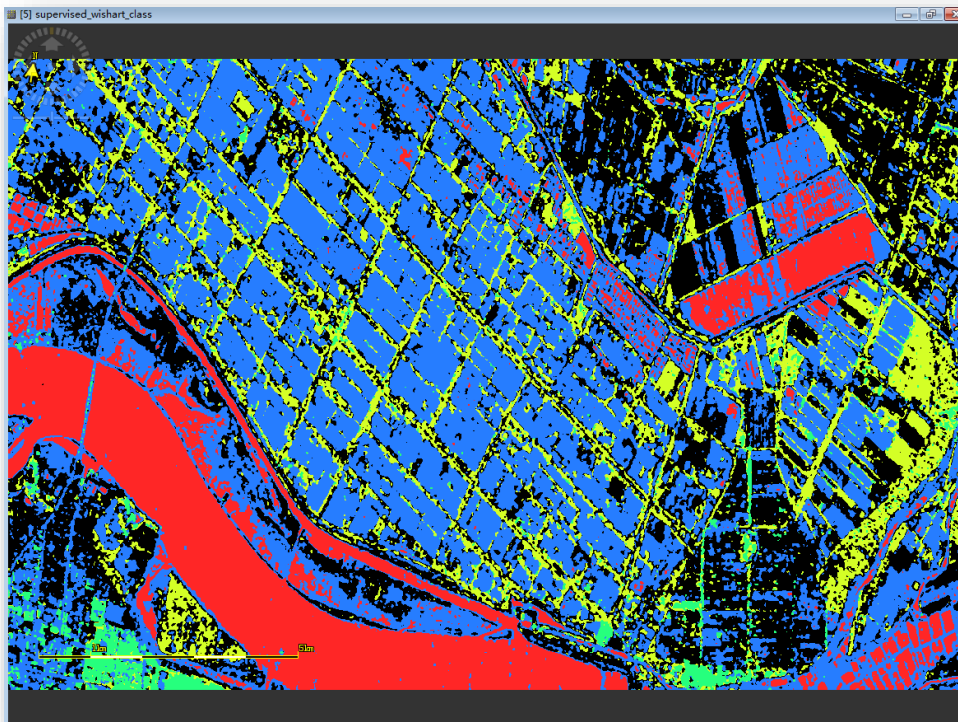
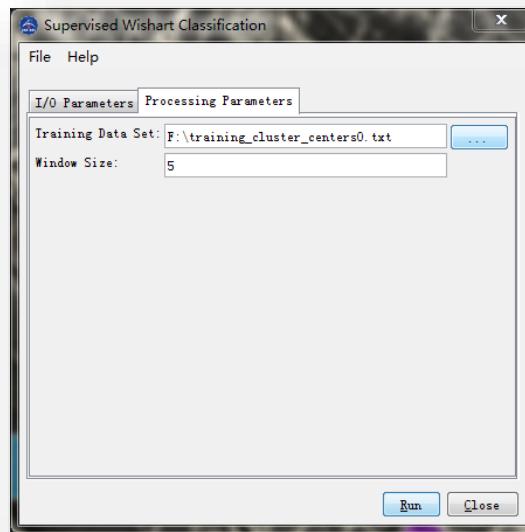




From the **SAR Processing** menu, go to the **Polarimetric** menu and select **Supervised Wishart Classification**.



In the **Processing Parameters** tab, select the training geometries txt file, window size 5x5. Press **Run** to begin processing.



Crops mapping with polarimetric SAR Data using GF-3 data

Objectives:

How to process GF-3 polarimetric products using PolSARpro 6.0

Polarimetric decompositions of GF-3 data for Crops identification

Crops mapping using unsuperivised polarimetric classification and supervised wishart classification

Data: The sample data used in this exercise was a subset from one GF-3 quad-pol image acquired on July 6, 2017

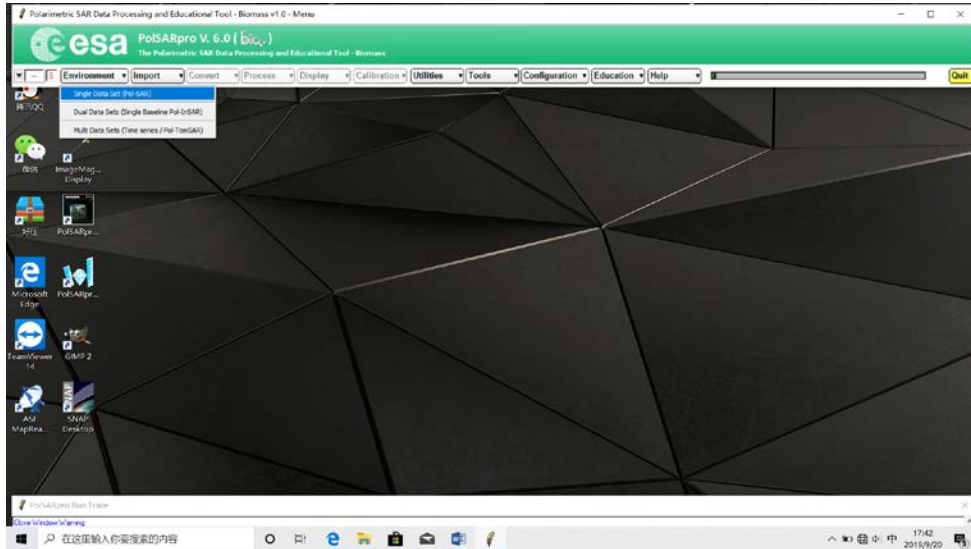
Product	Single Look Complex (SLC)
Incidence angle mid swath	41.18
Nominal Resolution	8
Location	Guangxi, China
Center Frequency	5.40

Software: PolSARpro 6.0

Tutorial:

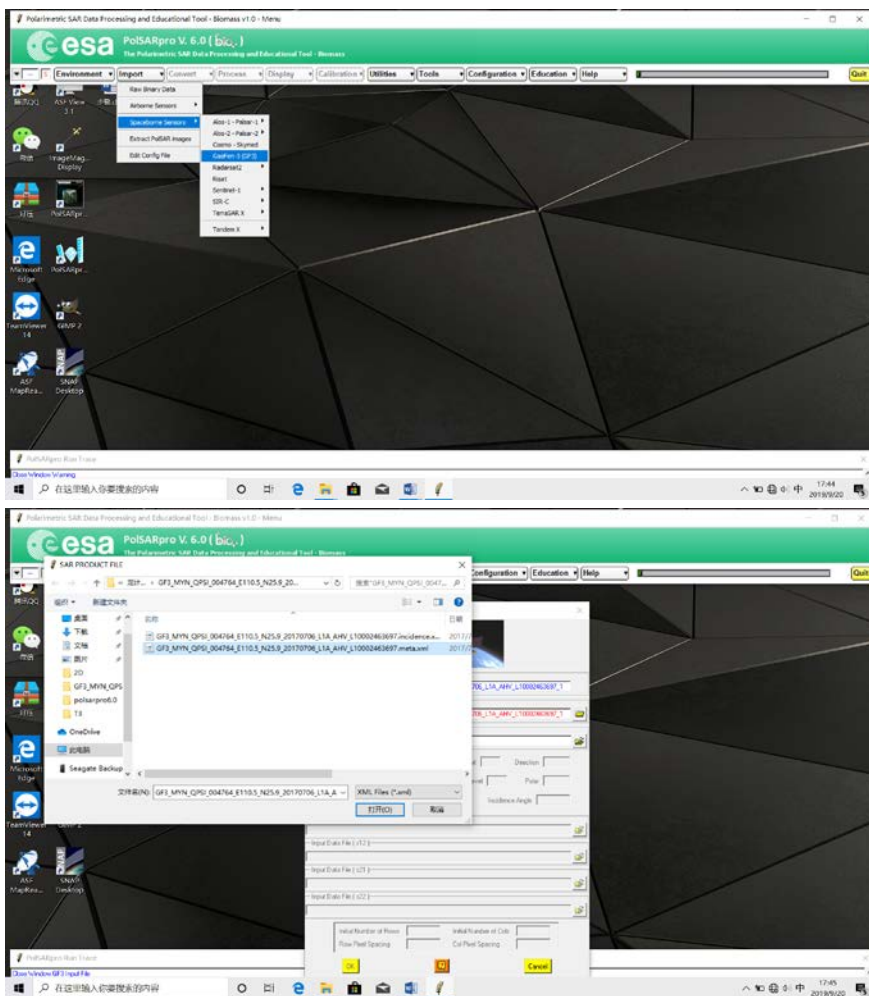
We choose the environment: Environment----Single Data Set (Pol-SAR), then choosing GF-3 data file folder.

Selected and click “Save&Exit”, Will pop up one Warning: Choose “No”.



import data: import----Spaceborne Sensors----GaoFen-3(GF-3)

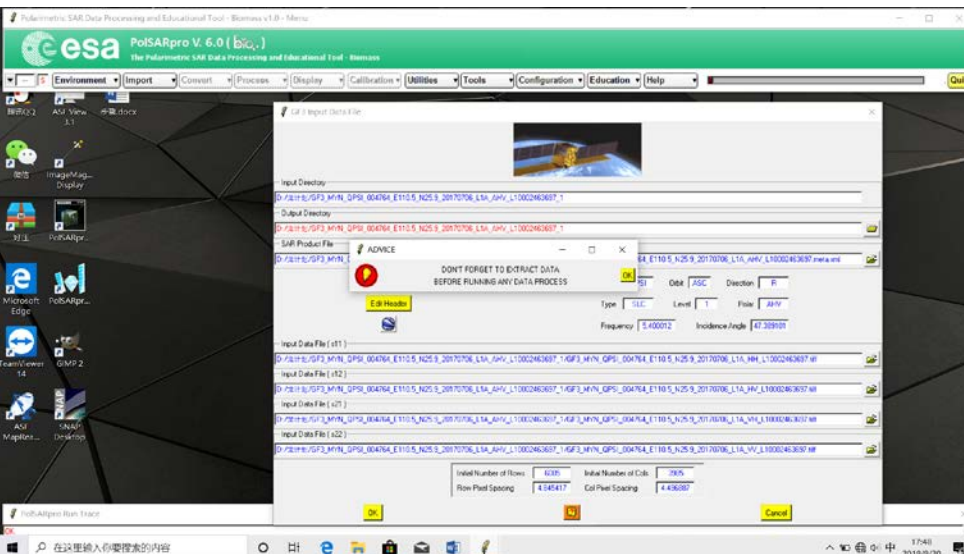
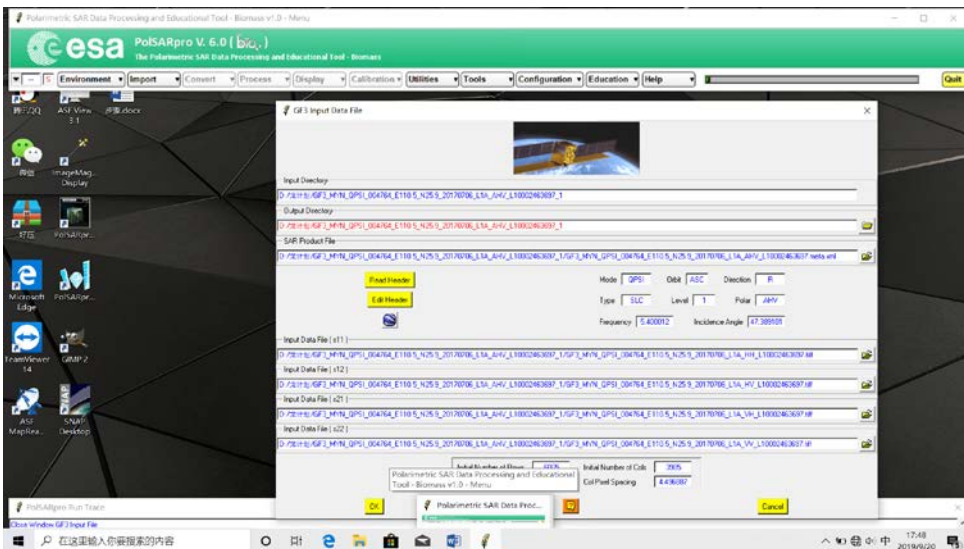
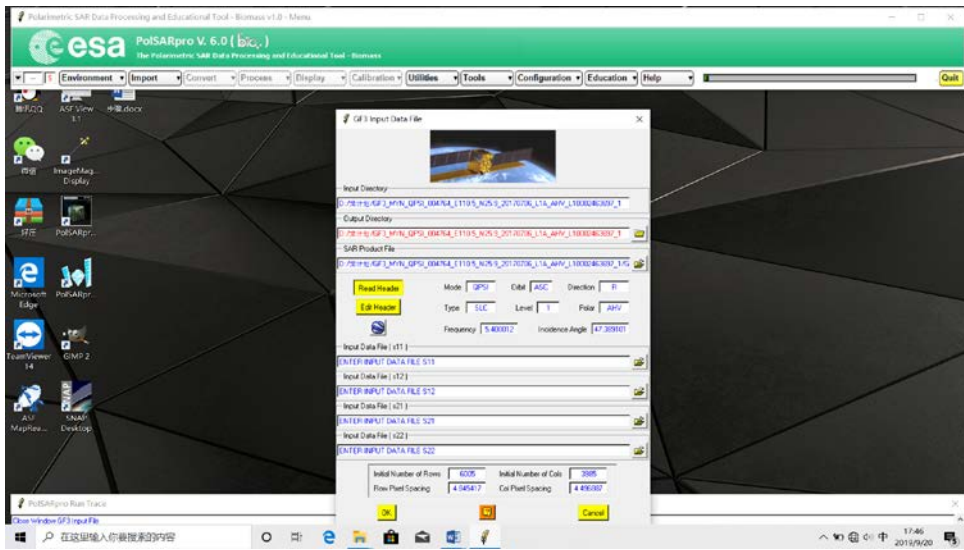
open file `GF3_MYN_QPSI_004764_E110.5_N25.9_20170706_L1A_AHV_L10002463697.meta.xml`



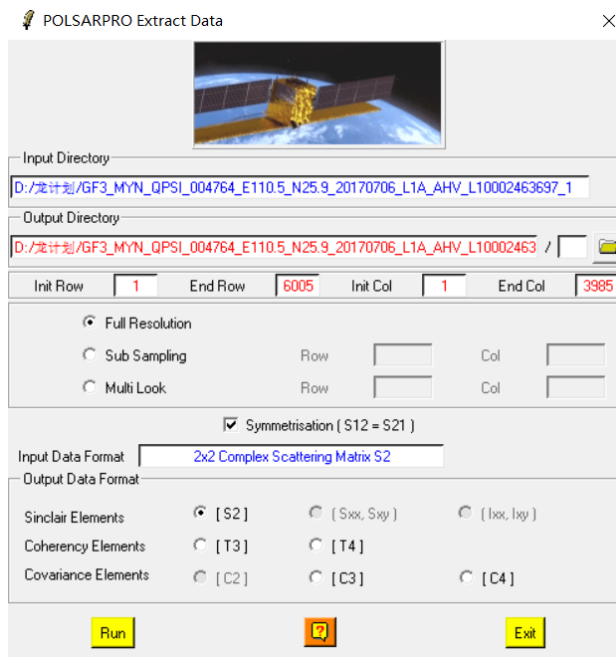
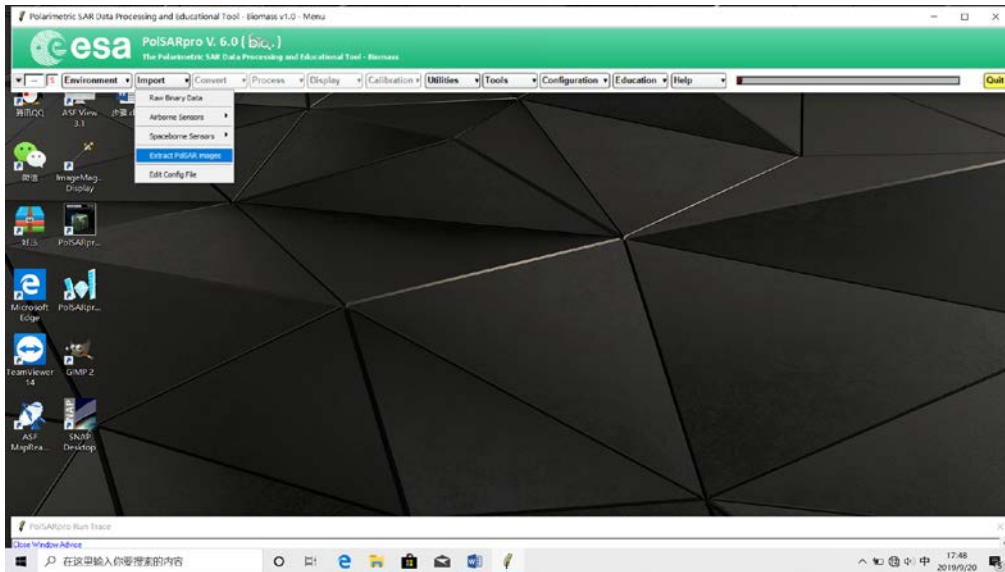
Read head file: Click

Read Header

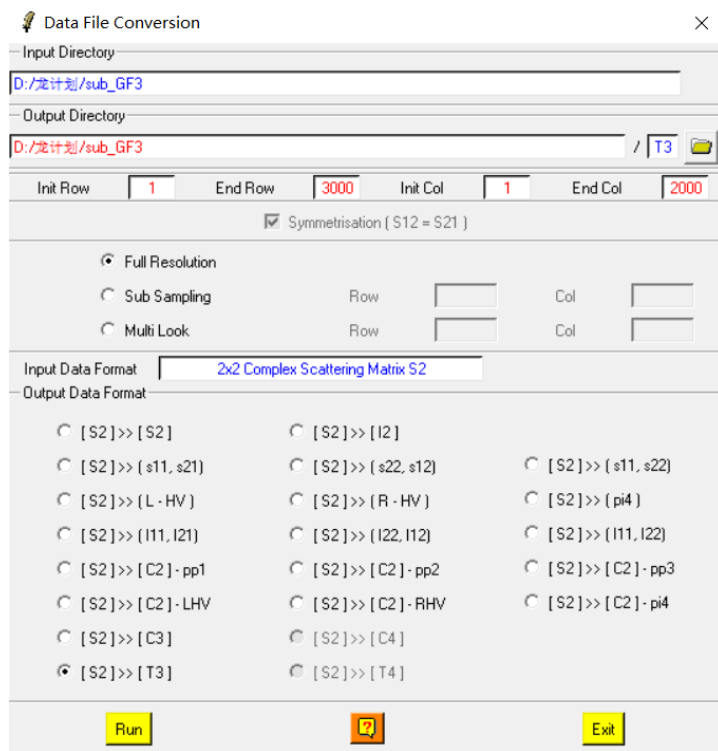
Import four polarizations S11\S12\S21\S22 corresponding to HH\HV\ VH\VV



Extract PolSAR images, and choose Full Resolution. out data format Choose Sinclair Element.

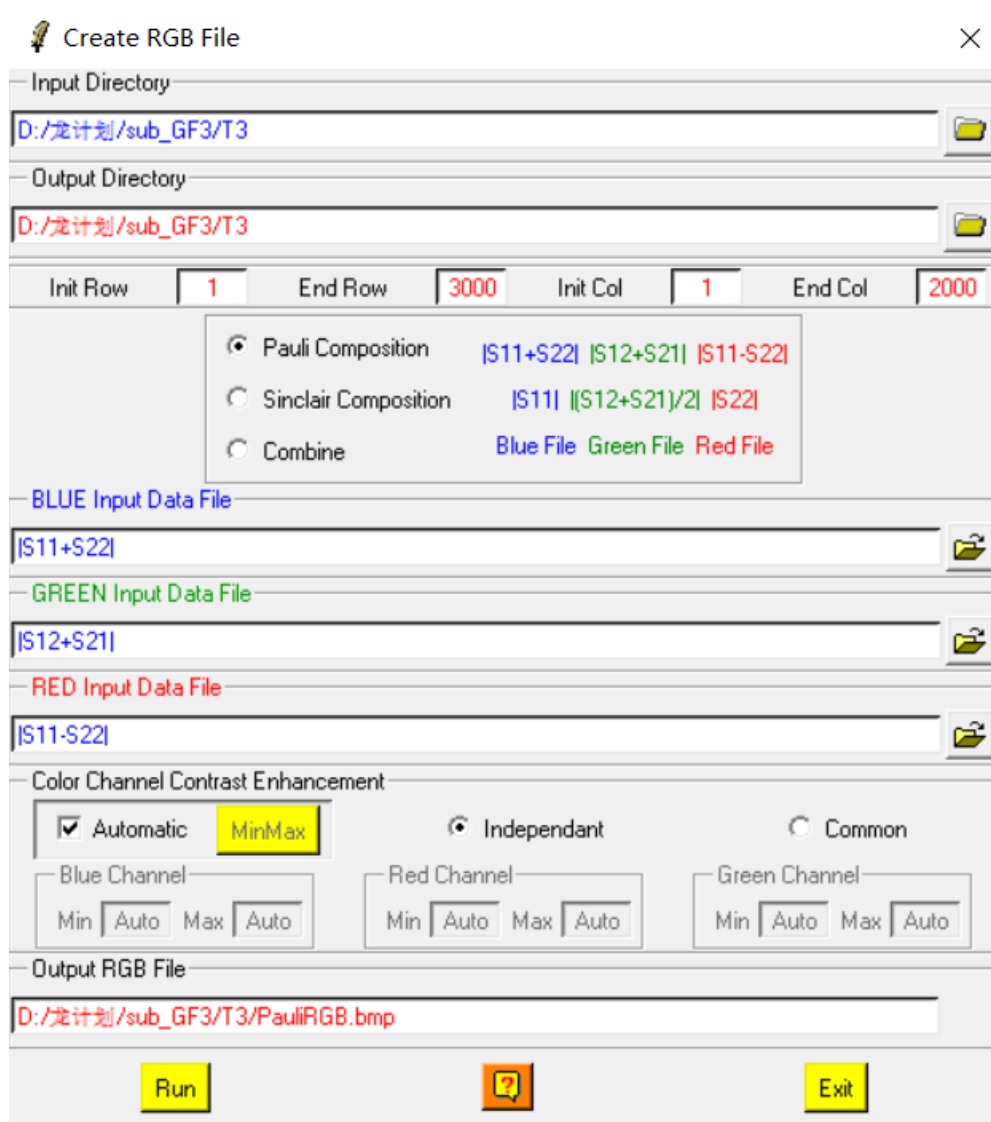


data conversion: Converting S2 to T3



Generate an RGB image for displaying a color composite image

Select: Display—>Create RGB File—>RGB Color Composition 1—>Run

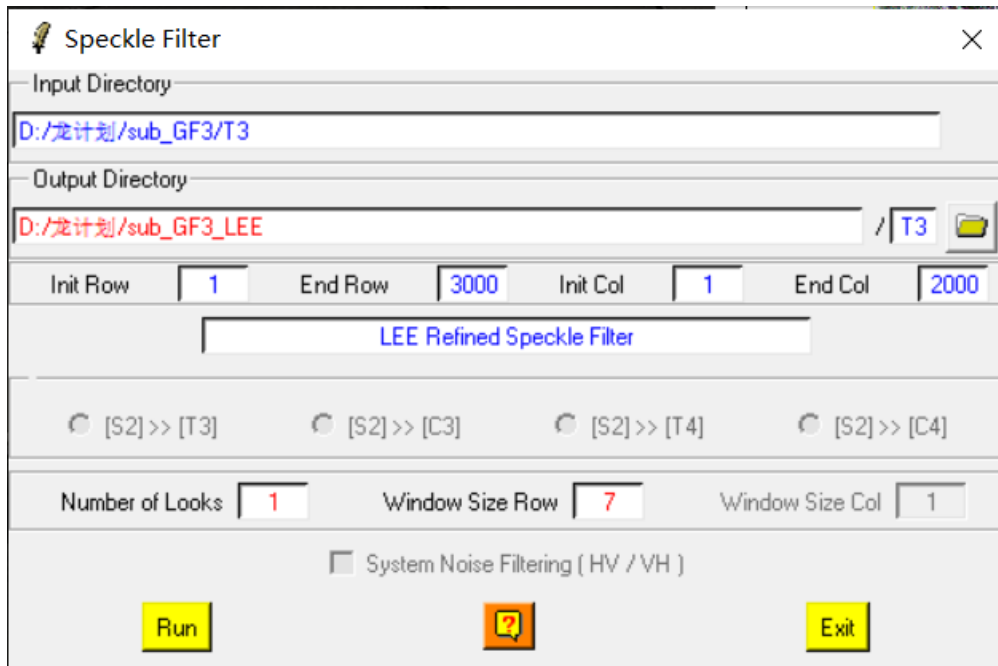


To clean up some of the speckle inherent in SAR images, you can apply a speckle filter. For full polarimetric data, speckle filters must take advantage of all bands and preserve the complex information.

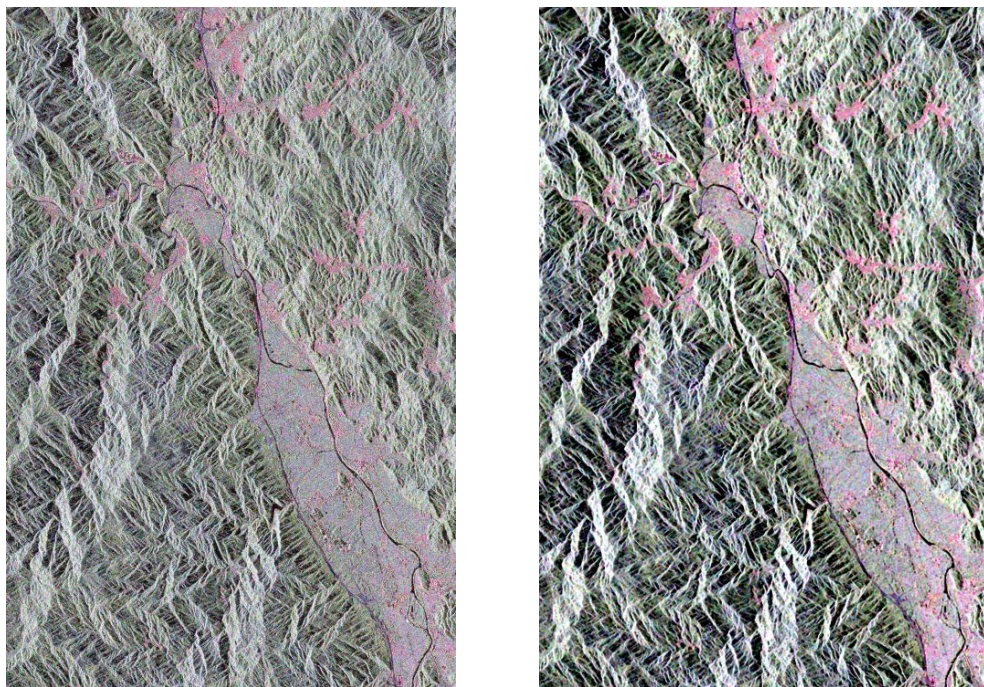
Next, the matrix T3 element is filtered.

Select: Process—>Polarimetric Speckle Filter—>Lee Refined Filter

Lee Refined Filter, the reason for choosing this filter is that it can retain the polarization information well under the influence of removing speckle filters.



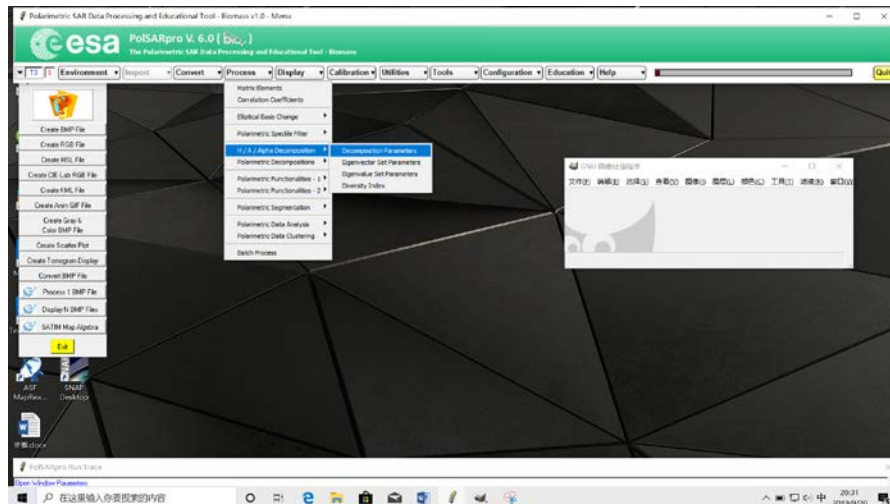
滤波前和滤波后的 RGB 合成图对比



Polarimetric Decompositions

Polarimetric decompositions allow the separation of different scattering contributions and can be used to extract information about the scattering process. From Process menu, elect Polarimetric Decomposition.

For example H/A/Alpha Polarimetric Decomposition



Data Processing: H / A / Alpha Decomposition Parameters

Input Directory
D:/龙计划/sub_GF3_LEE/T3

Output Directory
D:/龙计划/sub_GF3_LEE / T3

Init Row 1 End Row 3000 Init Col 1 End Col 2000

Alpha, Beta, Delta, Gamma, Lambda BMP

Lambda BMP

Alpha BMP

Entropy (H) BMP

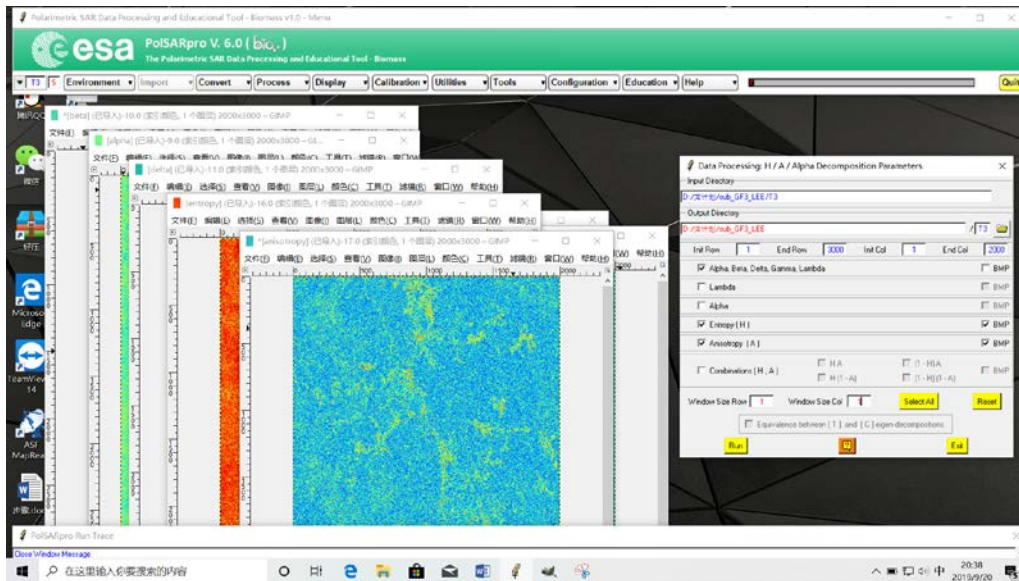
Anisotropy (A) BMP

Combinations (H , A) H A (1 - H) A BMP

H (1 - A) (1 - H) (1 - A)

Window Size Row 1 Window Size Col 1

Equivalence between [T] and [C] eigen-decompositions.



Unsupervised polarimetric classification

H/A/Alpha Classification: choose Process → Polarimetric Segmentation → H/A/Alpha Classification

